

INFORMATION DISCLOSURE CITATION

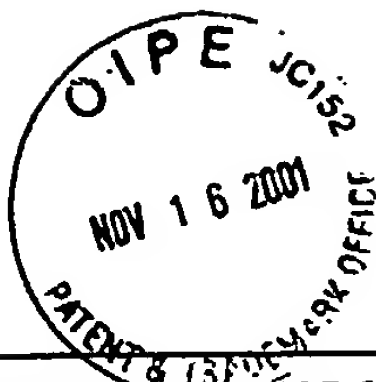
Atty. Docket No.	7414.0025-01	Appln. No.	09/911,903
Applicant	Hugh J. Pasika et al.		
Filing Date	July 23, 2001	Group:	8661 1631

U.S. PATENT DOCUMENTS							
Examiner Initial*	Document Number	Issue Date	Name	Class	Sub Class	Filing Date if appropriate	
cm	5,580,728	Dec 3, 1996	Perlin				
	6,019,896	Feb 1, 2000	Mansfield et al.				

FOREIGN PATENT DOCUMENTS							
	Document Number	Publication Date	Country	Class	Sub Class	Translation Yes or No	
ca	WO 99/53423	Oct 21, 1999	PCT				

OTHER DOCUMENTS (Including Author, Title, Date, Pertinent Pages, Etc.)	
ca	Ng, See-Kiong, "Automating Computational Molecular Genetics: Solving the Microsatellite Genotyping Problem,," School of Computer Science, Carnegie Mellon University, Pittsburgh, PA 15213, Jan 23, 1998, submitted in partial fulfillment of the requirements for the degree of Doctor of Philosophy, pages 107-150.
	Pálsson, Birgir, et al., "Using Quality Measures to Facilitate Allele Calling in High-Throughput Genotyping," <u>Genome Research</u> , Vol. 9, No. 10, Oct 1999, pages 1002-1012.
	Perlin, Mark W., "An Expert System for Scoring DNA Database Profiles," Genetic Identity Conference Proceedings, 11 th International Symposium on Human Identification - 2000, Oct 7-14, 2000, Biloxi, MS, 30 pages.

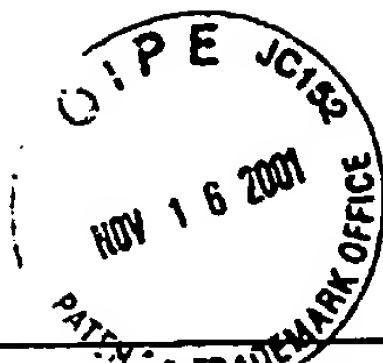
Examiner	<i>altur</i>	Date Considered	4/16/04
*Examiner:	Initial if reference considered, whether or not citation is in conformance with MPEP 609; draw line through citation if not in conformance and not considered. Include copy of this form with next communication to applicant.		
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U.S. PATENT DOCUMENTS							
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COL	ABI PRISM® Genotyper® 2.5 User's Manual, PE Biosystems, 1998, Foster City, CA.
COL	"Genetic Profiler Software," [online], [Retrieved on 5/15/2001]. Retrieved from http://www.mdyn.com/products/genotyping/software.htm , 3 pages.
	"Genotyping: High-throughput Genotyping," 5/15/2001, http://www.apbiotech.com/product/publication/lisn/lisn4/lisn4-1.html , 4 pages.
	Hoffman, E.P. et al., 1993, "Fully Automated PCR-based Disease Gene Mapping," <u>Am. J. Hum. Genet.</u> , 53(3 Supplement):Abstract 1172.
	Lancia, G. et al., 1998, "Genotyping of Pooled Microsatellite Markers by Combinatorial Optimization Techniques," <u>Discrete Applied Mathematics</u> , 88:291-314.
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	Ng, S.-K. et al., 1996, "Fully Automated Microsatellite Genotyping: Computer Software," <u>Am. J. Hum. Genet.</u> , 59(4 Supplement):Abstract 1319.
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	Perlin, M. W. et al., 1994, "Toward Fully Automated Genotyping: Allele Assignment, Pedigree Construction, Phase Determination, and Recombination Detection in Duchenne Muscular Dystrophy," <u>Am. J. Hum. Genet.</u> , 55:777-787.

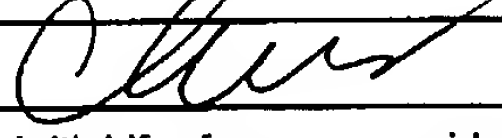


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	"Press Release: Amersham Pharmacia Biotech Unveils New Second Generation Software for MegaBACE 1000 DNA Sequencer," [online], [Retrieved on 5/15/2001], Retrieved from http://www.apbiotech.com/product/product_news/mega1000.htm , 2 pages.
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